RECEIVED

OCT 1 0 2002

TECH CENTER 1600/2900



1600

Page 1 of 7

RAW SEQUENCE LISTING DATE: 10/01/2002 PATENT APPLICATION: US/09/441,411 TIME: 15:46:17

Input Set : D:\409.app.txt

	<110> APPLICANT: Scholler, Nathalie B.	 -
5	•	ENTERED
	Hellstrom, Ingegerd	ENIL
7	,	
	<120> TITLE OF INVENTION: SURFACE RECEPTOR ANTIGEN	VACCINES
	<130> FILE REFERENCE: 730033.409	
	<140> CURRENT APPLICATION NUMBER: US 09/441,411	
	<141> CURRENT FILING DATE: 1999-11-16	
	<160> NUMBER OF SEQ ID NOS: 26	
	<170> SOFTWARE: FastSEQ for Windows Version 4.0	
	<210> SEQ ID NO 1	
	<211> LENGTH: 29	
	<212> TYPE: DNA	
24	<213> ORGANISM: Artificial Sequence	
	<220> FEATURE:	
	<223> OTHER INFORMATION: PCR primer	
	<400> SEQUENCE: 1	
30	ctaagcttat ggcttgcaat tgtcagttg	29
	<210> SEQ ID NO: 2	
	<211> LENGTH: 29	
34	<212> TYPE: DNA	
35	<213> ORGANISM: Artificial Sequence	
	<220> FEATURE:	
38	<223> OTHER INFORMATION: PCR primer	
40	<400> SEQUENCE: 2	
41	gtatcgatct aaaggaagac ggtctgttc	29
43	<210> SEQ ID NO: 3	
44	<211> LENGTH: 27	
45	<212> TYPE: DNA	
46	<213> ORGANISM: Artificial Sequence	
48	<220> FEATURE:	
49	<223> OTHER INFORMATION: PCR primer	
51	<400> SEQUENCE: 3	
52	cgaagcttgt tccagaactt acggaag	27
54	<210> SEQ ID NO: 4	
55	<211> LENGTH: 26	
56	<212> TYPE: DNA	
57	<213> ORGANISM: Artificial Sequence	
59	<220> FEATURE:	
60	<223> OTHER INFORMATION: PCR primer	
	<400> SEQUENCE: 4	
63	cgatcgatct ttcctcaggc tctcac	26
65	<210> SEQ ID NO: 5	

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/441,411 TIME: 15:46:17

DATE: 10/01/2002

Input Set : D:\409.app.txt

66 <211> LENGTH: 4473	
67 <212> TYPE: DNA	
68 <213> ORGANISM: Homo sapiens	
70 <400> SEQUENCE: 5	
71 aaggggaggt aaccetggee cetttggteg gggeeeeggg cageegegeg eeeetteeca	60
72 eggggeeett tactgegeeg egegeeegge ceccaecect egeageacee egegeeege	120
73 gocotoccag cogggtocag coggagocat ggggooggag cogcagtgag caccatggag	180
74 ctggcggcct tgtgccgctg ggggctcctc ctcgccctct tgccccccgg agccgcgagc	240
75 acceaagtgt geaceggeac agacatgaag etgeggetee etgecagtee egagacecae	300
76 ctggacatgc teegecacet etaceaggge tgecaggtgg tgeagggaaa cetggaacte	360
77 acctacetge ceaccaatge cageetgtee tteetgeagg atatecagga ggtgeaggge	420
78 tacgtgetca tegetcacaa ecaagtgagg caggteecae tgeagagget geggattgtg	480
78 cgaggcaccc agetetttga ggacaactat geeetggeeg tgetagacaa tggagacceg	540
80 ctgaacaata ccaccctgt cacaggggcc tccccaggag gcctgcggga gctgcagctt	600
81 cgaageetca cagagatett gaaaggaggg gtettgatee ageggaacee eeagetetge	660
82 taccaggaca cgattttgtg gaaggacatc ttccacaaga acaaccagct ggctctcaca	720
83 ctgatagaca ccaaccgctc tegggeetge cacccetgtt etecgatgtg taagggetee	780
84 cgctgctggg gagagagttc tgaggattgt cagagcctga cgcgcactgt ctgtgccggt	840
85 ggctgtgccc gctgcaaggg gccactgccc actgactgct gccatgagca gtgtgctgcc	900
	960
	1020
87 atotytyago tycactycec ageoctygic acctacada cayacacytt tydytecaty 88 occaatocog aggycogyta tacattogyc gocayctyty tyactycoty tocctacaac	1080
89 tacetteta eggacgtggg atcetgeace etegtetgee ecetgeacaa ecaagaggtg	1140
90 acagcagagg atggaacaca gcggtgtgag aagtgcagca agccctgtgc ccgagtgtgc	1200
91 tatggtetgg geatggagea ettgegagag gtgagggeag ttaccagtge caatatecag	1260
92 gagtttgctg gctgcaagaa gatctttggg agcctggcat ttctgccgga gagctttgat	1320
93 ggggacccag cetecaacae tgeceegete cagecagage agetecaagt gtttgagaet	1380
94 ctggaagaga tcacaggtta cctatacatc tcagcatgge cggacagcct gcctgacctc	1440
95 agogtottoc agaacetgca agtaateegg ggacgaatte tgcacaatgg egectacteg	1500
96 ctgaccctgc aagggctggg catcagetgg ctggggctgc gctcactgag ggaactgggc	1560
97 agtggactgg coctcatoca coataacaco cacototgot togtgoacac ggtgcootgg	1620
98 gaccagetet tteggaacee geaceaaget etgeteeaca etgecaaceg gecagaggae	1680
99 gagtgtgtgg gcgagggcct ggcctgccac cagctgtgcg cccgagggca ctgctggggt	1740
100 ccagggccca cccagtgtgt caactgcagc cagttccttc ggggccagga gtgcgtggag	1800
101 gaatgoogag tactgoaggg gctccccagg gagtatgtga atgccaggca ctgtttgccg	1860
102 tgccaccotg agtgtcagcc ccagaatggc tcagtgacct gttttggacc ggaggctgac	1920
103 cagtgtgtgg cotgtgccca ctataaggac cotcocttot gogtggcccg otgccccage	1980
104 ggtgtgaaac ctgacctctc ctacatgccc atctggaagt ttccagatga ggagggcgca	2040
105 tgccagcett gccccatcaa ctgcacccac tcctgtgtgg acctggatga caagggctge	2100
106 cccgccgage agagagecag ccctctgacg tecateatet ctgcggtggt tggcattetg	1160
107 ctggtcgtgg tcttgggggt ggtctttggg atcctcatca agcgacggca gcagaagate	2220
108 cggaagtaca cgatgcggag actgctgcag gaaacggagc tggtggagcc gctgacacct	2280
109 ageggagega tgcccaacca ggegeagatg eggateetga aagagaegga getgaggaag	2340
110 gtgaaggtgc ttggatctgg cgcttttggc acagtctaca agggcatctg gatccctgat	2400
111 ggggagaatg tgaaaattcc agtggccatc aaagtgttga gggaaaacac atcccccaaa	2460
112 gccaacaaag aaatettaga egaageatae gtgatggetg gtgtgggete eccatatgte	2520
113 tecegeette tgggeatetg cetgacatee acggtgeage tggtgacaca gettatgee	2580
114 tatggetgee tettagacea tgteegggaa aacegeggae geetgggete ecaggacetg	2640
115 ctgaactggt gtatgcagat tgccaagggg atgagctace tggaggatgt geggetegta	2700
IID digaadiggi glaigdagai iyodaayyyy alyayddadd lyydyyddyd gdyddiga	2

RAW SEQUENCE LISTING

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/441,411

DATE: 10/01/2002
TIME: 15:46:18

Input Set : D:\409.app.txt

116	cacagggact	tggccgctcg	gaacgtg	rctg	gtca	agag	jtc	ccaa	ccat	gt (caaaa	ttaca	2760
117	gacttcgggc	tggctcggct	gctggac	att	gaco	gagac	cag	agta	ccat	gc a	agatg	ggggc	2820
118	aaggtgccca	tcaagtggat	ggegetg	gag	teca	attct	CCC	gccg	gcgg	tt (caccc	accag	2880
119	agtgatgtgt	ggagttatgg	tgtgact	gtg	tggg	gaget	.ga	tgac	tttt	gg (ggcca	aacct	2940
120	tacqatqqqa	teccageeeg	ggagato	ecct	gaco	etget	.gg	aaaa	gggg	ga 🔻	gcggc	tgccc	3000
121	cageceecca	tctqcaccat	tgatgto	ctac	atga	atcat	_gg	tcaa	atgt	tg (gatga	ttgac	3060
122	totgaatgto	ggccaagatt	ccgggag	ıttg	gtgt	cctga	aat	tctc	ccgc	at (ggcca	gggac	3120
123	ccccaqcqct	ttgtggtcat	ccagaat	gag	gact	ttggg	jec	cage	cagt	CC	cttgg	acagc	3180
124	accttctacc	gctcactgct	ggaggad	gat	gaca	atggg	ggg	acct	ggtg	ga	tgctg	aggag	3240
125	tatctqqtac	cccaqcaqqq	cttcttc	etgt	ccag	gacco	ctg	CCCC	gggc	gc :	tgggg	gcatg	3300
126	qtccaccaca	ggcaccgcag	ctcatct	acc	agga	agtgg	geg	gtgg	ggac	ct	gacac	taggg	3360
127	ctggagccct	ctgaagagga	ggccccc	cagg	tete	ccact	-gg	cacc	ctcc	ga	agggg	ctggc	3420
128	tocgatgtat	ttgatggtga	cctggga	atg	gggg	gcago	cca	aggg	gctg	ca	aagcc	tcccc	3480
129	acacatgacc	ccagccctct	acagege	jtac	agto	gagga	acc	ccac	agta	CC	cctgc	cctct	3540
130	gagactgatg	gctacgttgc	cccctc	jacc	tgca	agcco	ccc	aged	tgaa	ta	tgtga	accag	3600
1.3.1	ccagatgttc	ggccccagcc	cccttcc	, dcc	cgag	gaggg	jcc	ctct	gcct	.gc	tgccc	gacct	3660
132	gctggtgcca	ctctqqaaaq	gcccaac	jact	ctct	tacco	cag	ggaa	gaat	gg ·	ggtcg	tcaaa	3720
133	gacgtttttg	cctttggggg	taccata	gag	aaco	edega	agt	actt	gaca	CC	ccagg	gagga	3780
134	getgeeete	agccccaccc	tectect	gcc	ttca	agcco	cag	cctt	cgac	aa	cctct	attac	3840
135	tgggaccagg	acccaccaga	acadada	get	ccad	ccca	gca	cctt	.caaa	gg	gacac	ctacg	3900
136	gcagagaacc	cagagtacct	gggtctg	ggac	gtgo	ccagi	tgt.	gaac	caga	ag	gccaa	gtccg	3960
137	gcagagaacc cagagtacct gggtctggac gtgccagtgt gaaccagaag gccaagtccg cagaagccct gatgtgtcct cagggagcag ggaaggcctg acttctgctg gcatcaagag										4020		
138	gtgggagggc	cctccgacca	cttccac	adad	aaco	ctgc	cat	gcca	ggaa	CC	tgtcc	taagg	4080
139	aaccttcctt	cctgcttgag	ttcccad	atq	gcto	ggāa	ggg	gtcc	agco	tc	gttgg	aagag	4140
140	gaacagcact	ggggagtett	tatagat	tct	gage	gecet	tqc	ccaa	tgag	ac	tctag	ggtcc	4200
1.11	gaacagcact ggggagtett tgtggattet gaggeeetge eeaatgagae tetagggtee agtggatgee acageecage ttggeeettt eetteeagat eetgggtaet gaaageetta										4260		
142	I adiqually acadeceade regardere entre ent											4320	
1.13	A UUUGGUCLUU CCCUGUUUUGU GUUGOGGOOO CUUGGGGGGGGG											4380	
1.1.4	j atticadadac tycocotyda acocagoace goodateer, njjiinji											4440	
	ttttttaaag					_	_						4473
	<210> SEQ		J	,,,,	,								
	<211> LENG												
	<212> TYPE												
	<213> ORGA		sapiens										
	<400> SEQU		- L										
153	Met Glu Le	u Ala Ala L	eu Cvs A	Arq 1	Trp (Gly :	Leu	Leu	Leu	Ala	Leu	Leu	
154		5	1	,		10					15		
155	Pro Pro Gl		er Thr (iln v	Val (Cvs '	Thr	Gly	Thr	Asp	Met	Lys	
156	FIO TIO GI	20			25	-1-		-		30		-	
157	Leu Arg Le	u Pro Ala S	er Pro (His 1	Leu	Asp	Met	Leu	Arg	His	
158				10					45				
150	Leu Tyr Gl				Gln (Glv i	Asn	Leu	Glu	Leu	Thr	Tyr	
160	50	n diy cyb d	55			O-1 .		60				•	
161	Leu Pro Th	r Acn Ala S		Ser 1	Phe '	Leu (Gln		Ile	Gln	Glu	Val	
$\frac{161}{162}$			0				75					80	
162	Gln Gly Ty			His A	Asn (Ara	Gln	Val	Pro		
164	GIU GIA IA	85	LU MILU I			90		9			95		
165	Gln Arg Le		al Ara (Glv ′			Leu	Phe	Glu	Asp		Tyr	
166	oin ary be	100	··- ·		105					110		-	
T 0 0		100											

RAW SEQUENCE LISTING DATE: 10/01/2002 PATENT APPLICATION: US/09/441,411 TIME: 15:46:18

Input Set : D:\409.app.txt

167	Ala	Leu		Val	Leu	Asp	Asn		Asp	Pro	Leu	Asn		Thr	Thr	Pro
168			115					120					125			
169	Val	Thr	Gly	Ala	Ser	Pro	Gly	Gly	Leu	Arg	Glu	Leu	Gln	Leu	Arg	Ser
170		130					135					140				
171	Leu	Thr	Glu	Ile	Leu	Lys	Gly	Gly	Val	Leu	Ile	Gln	Arg	Asn	Pro	Gln
	145					150					155					160
173	Leu	Cys	Tyr	Gln	Asp	Thr	Ile	Leu	Trp	Lys	Asp	Ile	Phe	His	Lys	Asn
174		•	-		165					170					175	
	Asn	Gln	Leu	Ala	Leu	Thr	Leu	Ile	Asp	Thr	Asn	Arg	Ser	Arg	Ala	Cys
176				180	-				185			_		190		_
	His	Pro	Cvs		Pro	Met	Cvs	Lvs	Glv	Ser	Arg	Cvs	Trp	Gly	Glu	Ser
178	1115	110	195	001			-1-	200	1		,	4	205	-		
	Ser	Glu		Cvs	G1 n	Ser	Len		Ara	Thr	Val	Cvs	Ala	Glv	Glv	Cvs
180	DCI	210	тор	010	QIII	001	215					220		1	- 1	-
	Ala		Cve	Lvc	Glv	Pro		Pro	Thr	Asp	Cvs		His	Glu	Gln	Cvs
	225	nry	Cys	цуз	Gry	230	пси	110	1	p	235	010		014	02	240
	Ala	λla	C111	Cvc	Thr		Dro	Lve	Hic	Ser		Cvs	I.eu	Δla	Cvs	
	Ala	Ата	СТУ	СУЗ	245	ату	FIU	цуз	1113	250	пор	Cys	пси	711 CI	255	LCu
184	His	Dha	3.00	ni a		C1	т1о	Cvc	Clu		Uic	Cvc	Dro	λla		Val
	HIS	Pile	ASII		261	GIY	iie	Cys	265	Leu	1113	Cys	110	270	пси	***
186	m 1	m	3	260	3	m b so	Dho	C1.		Mot	Dro	λan	Dro		Clu	λνα
	Thr	Tyr		Thr	ASP	THE	Pne		ser	Met	PIO	ASII	285	Gru	Gry	Arg
188	_		275	a1	31.	a		280	m la sa	7 l a	Crra	Dro		Nan	Фтт∞	Tou
	Tyr		Phe	GTÄ	Ата	ser		vaı	Thr	Ата	Cys		тут	ASII	1 y 1	ьеu
190	_	290			-1	_	295	m1	.	17 - 1	Q	300	T	TI	3 an	C1.5
	Ser	Thr	Asp	Val	GLY		Cys	Thr	Leu	vaı		Pro	Leu	HIS	ASII	
	305	_			- 1	310	- 1	 1	~ 3		315	a1	T +	G	C	320
	Glu	Val	Thr	Ala		Asp	GIY	Thr	GIn		Cys	GIU	Lys	Cys		Lys
194					325	_	_	- 1	_	330		~ 1		.	335	a1
	Pro	Cys	Ala		Val	Cys	Tyr	GLY		GTA	Met	G1u	HlS		Arg	GIU
196				340					345	_		_,		350	_	_
	Val	Arg		Val	Thr	Ser	Ala		Ile	Gln	Glu	Phe		GLY	Cys	Lys
198			355					360					365			_
199	Lys	Ile	Phe	Gly	Ser	Leu		Phe	Leu	Pro	Glu		Phe	Asp	Gly	Asp
200		370					375					380		_		
201	Pro	Ala	Ser	Asn	Thr	Ala	Pro	Leu	Gln	Pro		Gln	Leu	Gln	Val	
-	385					390					395					400
203	Glu	Thr	Leu	Glu	Glu	Ile	Thr	Gly	Tyr		Tyr	Ile	Ser	Ala		Pro
204					405					410					415	
205	Asp	Ser	Leu	Pro	Asp	Leu	Ser	Val	Phe	Gln	Asn	Leu	Gln	Val	Ile	Arg
206				420					425					430		
207	Gly	Arg	Ile	Leu	His	Asn	Gly	Ala	Tyr	Ser	Leu	Thr	Leu	Gln	Gly	Leu
208			435					440					445			
209	Gly	Ile	Ser	Trp	Leu	Gly	Leu	Arg	Ser	Leu	Arg	Glu	Leu	Gly	Ser	Gly
210		450					455					460				
	Leu	Ala	Leu	Ile	His	His	Asn	Thr	His	Leu	Cys	Phe	Val	His	Thr	Val
	465					470					475					480
213	Pro	Trp	Asp	Gln	Leu	Phe	Arg	Asn	Pro	His	Gln	Ala	Leu	Leu	His	Thr
214		1	_		485					490					495	
	Ala	Asn	Ara	Pro		Asp	Glu	Cys	Val	Gly	Glu	Gly	Leu	Ala	Cys	His
			- 9					4 -	_	-		-			-	

RAW SEQUENCE LISTING DATE: 10/01/2002 PATENT APPLICATION: US/09/441,411 TIME: 15:46:18

Input Set : D:\409.app.txt

216				500					505					510		
217	Gln	Leu	Cys	Ala	Arg	Gly	His	Cys	Trp	Gly	Pro	Gly	Pro	Thr	Gln	Cys
218			515					520				_	525			_
	Val		Cys	Ser	Gln	Phe	Leu	Arg	Gly	Gln	Glu	Cys	Val	Glu	Glu	Cys
220		530					535					540				
			Leu	Gln	Gly	Leu	Pro	Arg	Glu	Tyr	Val	Asn	Ala	Arg	His	Cys
	545					550					555					560
	Leu	Pro	Cys	His	Pro	Glu	Cys	Gln	Pro	Gln	Asn	Gly	Ser	Val	Thr	Cys
224					565					570					575	
	Phe	Gly	Pro		Ala	Asp	Gln	Cys		Ala	Cys	Ala	His		Lys	Asp
226	_	_	_,	580					585					590		
	Pro	Pro		Cys	Val	Ala	Arg		Pro	Ser	Gly	Val		Pro	Asp	Leu
228	0	m	595	D	71.	m		600		_	a 1	a 3	605	- 1	_	- 1
	ser	610	мес	Pro	Ile	Trb		Pne	Pro	Asp	GIU		GLY	Ala	Cys	GIn
230	Dro		Dro	т10	Nan	Crra	615	II i o	Com	Crra	17-1	620	T 0.11	3	N	T
	625	Cys	PIO	116	Asn	630	1111	нтъ	ser	Cys	635	ASP	Leu	ASP	Asp	Lys 640
		Cvs	Pro	Δla	Glu		λησ	λla	Sar	Dro		Thr	Cor	T10	Tla	
234	Ory	Cys	110	niu	645	GIII	пту	AIG	261	650	пеа	1111	per	TIE	655	261
	Ala	Val	Va l	Glv	Ile	Leu	Leu	Val	Va 1		T.eu	Glv	Va 1	Val		Glv
236				660		200	DCu	, 41	665	, 41	нса	OI,	, u i	670	1 110	GLY
	Ile	Leu	Ile		Arg	Arq	Gln	Gln		Ile	Ara	Lvs	Tvr		Met.	Ara
238			675	1	,	,		680	1 -		5	-1-	685			5
239	Arg	Leu	Leu	Gln	Glu	Thr	Glu	Leu	Val	Glu	Pro	Leu	Thr	Pro	Ser	Gly
240	_	690					695					700				•
41	Ala	Met	Pro	Asn	Gln	Ala	${\tt Gln}$	Met	Arg	Ile	Leu	Lys	Glu	Thr	Glu	Leu
	705					710					715					720
	Arg	Lys	Val	Lys	Val	Leu	Gly	Ser	Gly	Ala	Phe	Gly	Thr	Val	Tyr	Lys
244					725					730					735	
	Gly	Ile	Trp		Pro	Asp	Gly	Glu		Val	Lys	Ile	Pro		Ala	Ile
246	_		_	740		_	_,	_	745					750	_	
	Lys	vaı		Arg	Glu	Asn	Thr		Pro	Lys	Ala	Asn	_	GIu	Ile	Leu
248	\ an	C1	755	m	1101	Mak	71.	760	17- 1	a 1	G	D	765	**- 7		_
250	ASP	770	Ald	171	Val	мес	775	GTÀ	vaı	GIA	ser		туг	vaı	ser	Arg
	Τωυ		C1v	Tlo	Cys	T OU		Cor	Thr	Wal	Cln	780	Wa 1	The	c1n	Τ Ο
252		цец	Gry	116	Суз	790	1111	ser	1111	val	795	Leu	vaı	1111	GIII	800
		Pro	Tvr	Glv	Cys		Leu	Asn	His	Va 1		Glu	Δen	Δrα	Clv	
254	1100	110	111	011	805	Leu	пса	пор	nii	810	1119	Olu	пып	лгу	815	Arg
	Leu	Glv	Ser	Gln	Asp	Leu	Leu	Asn	Trp		Met	Gln	Tle	Ala		Glv
256		- 1		820	1				825	010		0111		830	270	011
	Met.	Ser	Tyr		Glu	Asp	Val	Arq		Val	His	Arq	Asp		Ala	Ala
258			835			•		840					845	_	_	
259	Arg	Asn	Val	Leu	Val	Lys	Ser	Pro	Asn	His	Val	Lys	Ile	Thr	Asp	Phe
260		850					855					860				
261	Gly	Leu	Ala	Arg	Leu	Leu	Asp	Ile	Asp	Glu	Thr	Glu	Tyr	His	Ala	Asp
262						870					875					880
	Gly	Gly	Lys	Val	Pro	Ile	Lys	Trp	Met		Leu	Glu	Ser	Ile		Arg
264					885					890					895	

VERIFICATION SUMMARYDATE: 10/01/2002PATENT APPLICATION: US/09/441,411TIME: 15:46:19

Input Set : D:\409.app.txt